The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number:

Source:

Date Processed by STIC:

ENTERED



IFW16

RAW SEQUENCE LISTING

3 <110> APPLICANT: Andrade-Gordon, Patricia

DATE: 11/17/2004 TIME: 12:29:19

PATENT APPLICATION: US/10/040,655A

Input Set : A:\ORT1566NP.Subst.Seq.List.txt Output Set: N:\CRF4\11172004\J040655A.raw

```
Darrow, Andrew
         Qi, Jenson
 7 <120> TITLE OF INVENTION: CLEANING COMPOSITIONS CONTAINING HUMAN SERINE PROTEASE T
 9 <130> FILE REFERENCE: ORT-1566
11 <140> CURRENT APPLICATION NUMBER: US 10/040,655A
12 <141> CURRENT FILING DATE: 2002-01-07
14 <150> PRIOR APPLICATION NUMBER: US 09/386,653
15 <151> PRIOR FILING DATE: 1999-08-31
17 <160> NUMBER OF SEQ ID NOS: 11
19 <170> SOFTWARE: PatentIn version 3.3
21 <210> SEQ ID NO: 1
22 <211> LENGTH: 1110
23 <212> TYPE: DNA
24 <213> ORGANISM: Human
26 <400> SEQUENCE: 1
27 gaccacggcc ctgcgcccca gccaggcctg aggacatgag gcggccggcg gcggtgccgc
                                                                        60
29 teetgetget getgtgtttt gggteteaga gggeeaagge ageaacagee tgtggtegee
                                                                       120
31 ccaggatgct gaaccgaatg gtgggcgggc aggacacgca ggagggcgag tggccctggc
                                                                       180
33 aagtcagcat ccagcgcaac ggaagccact tctgcggggg cagcctcatc gcggagcagt
                                                                       240
35 gggtcctgac ggctgcgcac tgcttccgca acacctctga gacgtccctg taccaggtcc
                                                                       300
37 tgctggggc aaggcagcta gtgcagccgg gaccacacgc tatgtatgcc cgggtgaggc
39 aggtggagag caaccccctg taccagggca cggcctccag cgctgacgtg gccctggtgg
                                                                       420
41 agetggagge accagtgeee tteaccaatt acateeteee egtgtgeetg cetgaeecet
                                                                       480
43 cggtgatctt tgagacgggc atgaactgct gggtcactgg ctggggcagc cccagtgagg
                                                                       540
45 aagaceteet geeegaaceg eggateetge agaaactege tgtgeeeate ategacacae
                                                                       600
47 ccaagtgcaa cctgctctac agcaaagaca ccgagtttgg ctaccaaccc aaaaccatca
                                                                       660
49 agaatgacat gctgtgcgcc ggcttcgagg agggcaagaa ggatgcctgc aagggcgact
                                                                       720
51 cgggcggccc cctggtgtgc ctcgtgggtc agtcgtggct gcaggcgggg gtgatcagct
                                                                       780
53 ggggtgaggg ctgtgcccgc cagaaccgcc caggtgtcta catccgtgtc accgcccacc
                                                                       840
55 acaactggat ccatcggatc atccccaaac tgcagttcca gccagcgagg ttgggcggcc
                                                                       900
57 agaagtgaga cccccggggc caggagcccc ttgagcagag ctctgcaccc agcctgcccg
                                                                       960
59 cccacaccat cetgetggte etcecagege tgetgttgca cetgtgagee ccaccagact
                                                                      1020
1080
63 ccaataaaaa cccagcctgt gtgccagctg
                                                                      1110
66 <210> SEQ ID NO: 2
67 <211> LENGTH: 20
68 <212> TYPE: DNA
69 <213> ORGANISM: Artificial
71 <220> FEATURE:
72 <223> OTHER INFORMATION: ProtT PCRTP-U primer
74 <400> SEQUENCE: 2
75 gccaggcctg aggacatgag
```

20

PATENT APPLICATION: US/10/040,655A

DATE: 11/17/2004 TIME: 12:29:19

```
78 <210> SEQ ID NO: 3
 79 <211> LENGTH: 20
 80 <212> TYPE: DNA
 81 <213> ORGANISM: Artificial
 83 <220> FEATURE:
 84 <223> OTHER INFORMATION: ProtT PCRTP-L
 86 <400> SEQUENCE: 3
 87 tgcgctggat gctgacttgc
                                                                            20
 90 <210> SEQ ID NO: 4
 91 <211> LENGTH: 40
 92 <212> TYPE: DNA
 93 <213> ORGANISM: Artificial
 95 <220> FEATURE:
 96 <223> OTHER INFORMATION: ProtT PCRTP-PP
 98 <400> SEQUENCE: 4
 99 ccaggatgct gaaccgaatg gtgggcgggc aggacacgca
                                                                            40
 102 <210> SEQ ID NO: 5
103 <211> LENGTH: 30
104 <212> TYPE: DNA
105 <213 > ORGANISM: Artificial
107 <220> FEATURE:
108 <223> OTHER INFORMATION: Prot T Xba-U
110 <400> SEQUENCE: 5
111 aggatctaga ggaggggag tggccctggc
                                                                            30
114 <210> SEQ ID NO: 6
115 <211> LENGTH: 30
116 <212> TYPE: DNA
117 <213> ORGANISM: Artificial
119 <220> FEATURE:
120 <223> OTHER INFORMATION: Prot T Xba-L
122 <400> SEQUENCE: 6
123 ggggtctaga cttctggccg cccaacctcg
                                                                            30
126 <210> SEQ ID NO: 7
127 <211> LENGTH: 290
128 <212> TYPE: PRT
129 <213> ORGANISM: Human
131 <400> SEQUENCE: 7
133 Met Arg Arg Pro Ala Ala Val Pro Leu Leu Leu Leu Cys Phe Gly
137 Ser Gln Arg Ala Lys Ala Ala Thr Ala Cys Gly Arg Pro Arg Met Leu
138
                                     25
141 Asn Arg Met Val Gly Gly Gln Asp Thr Gln Glu Gly Glu Trp Pro Trp
142
            35
145 Gln Val Ser Ile Gln Arg Asn Gly Ser His Phe Cys Gly Gly Ser Leu
149 Ile Ala Glu Gln Trp Val Leu Thr Ala Ala His Cys Phe Arg Asn Thr
                                             75
153 Ser Glu Thr Ser Leu Tyr Gln Val Leu Leu Gly Ala Arg Gln Leu Val
154
                    85
```

PATENT APPLICATION: US/10/040,655A

DATE: 11/17/2004 TIME: 12:29:19

4				_													
15	7 Gln	Pro	GIY		His	Ala	Met	Tyr	Ala	Arg	Val	Arg	Gln	Val	Glu	Ser	
158				100					105					110			
161	. Asn	Pro	Leu	Tyr	Gln	Gly	Thr	Ala	Ser	Ser	Ala	Asp	Val	Ala	Len	Val	
162			115			-		120					125		Lea	val	
165	Glu	Leu	Glu	Ala	Pro	Val	Pro		Thr	Agn	Тиг	тла	LOU	Dro	570 T	<i></i>	
166		130			110	vai	135	LIIC	1111	Veri	TAT		ьeu	PIO	vai	Cys	
			7 ~~	Dwo	C	**- 7		D1	~7	1	~-7	140	_				
170	Leu	PIO	Asp	PIO	ser		тте	Pne	GIU	Thr		Met	Asn	Cys	Trp	Val	
	145	~ 7		~-7	_	150		_	_		155					160	
173	Thr	GLY	Trp	Gly		Pro	Ser	Glu	Glu	Asp	Leu	Leu	Pro	${ t Glu}$	Pro	Arg	
174					165					170					175		
177	Ile	Leu	Gln	Lys	Leu	Ala	Val	Pro	Ile	Ile	Asp	Thr	Pro	Lys	Cys	Asn	
178				180					185		_			190	-		
181	Leu	Leu	Tyr	Ser	Lys	Asp	Thr	Glu	Phe	Glv	Tvr	Gln	Pro	Lvs	Thr	Tla	
182			195		-	-		200		1	-1-		205	175		110	
185	Lys	Asn		Met	Leu	Cvs	Δla		Dho	Glu	Clu	C1		T	7 ~~	77-	
186	-1-	210				Cyb	215	O _T y	FIIC	Giu	Gru		пур	ьys	Asp	Ala	
			G1.	7 an	Cox	a 1		D	.	TT - 7	~	220					
100	Cys	пуъ	СТУ	Asp	ser		GIA	PIO	ьeu	vaı		Leu	Val	GLY	GIn	Ser	
	225	-	~ 7			230					235					240	
193	${\tt Trp}$	ьeu	GIn	Ala		Val	Ile	Ser	Trp	Gly	Glu	Gly	Cys	Ala	Arg	Gln	
194					245					250					255		
197	Asn	Arg	Pro	Gly	Val	Tyr	Ile	Arg	Val	Thr	Ala	His	His	Asn	Trp	Ile	
198				260					265					270			
201	His	Arg	Ile	Ile	Pro	Lys	Leu	Gln	Phe	Gln	Pro	Ala	Ara	Leu	Glv	Glv	
202			275			-		280			•		285		017	017	
205	Gln	Lvs											200				
206		290															
	<210		:O TI	NO.	B												
	<211> LENGTH: 1130 <212> TYPE: DNA																
	<pre><212> TIFE: DNA </pre> <pre><213> ORGANISM: Artificial</pre>																
	<pre><<220> FEATURE:</pre>																
	5 <223> OTHER INFORMATION: PFEK-PROTT-HIS fusion protein 7 <400> SEQUENCE: 8																
218	gaat	tcac	ca c	catg	gaca	g ca	aagg	ttcg	tcg	caga	aat	cccg	cctg	ct c	ctgc	tgctg	60
220	gtgg	tgtc	aa a	itcta	ctct	t gt	gcca	gggt	gtg	gtct	ccg	acta	caag	ga c	gacg	acgac	120
222	gtgg	acgc	gg c	cgct	cttg	c tg	CCCC	cttt	gat	gatg	atg	acaa	gatc	gt t	qqqq	qctat	180
224	gctc	taga	gg a	ıgggc	gagt	g gc	cctg	gcaa	gtc	agca	tcc	agcg	caac	- qq a	agcc	acttc	240
226	tgcg	9999	ca g	racta	atcg	c gg	agca	gtgg	gtc	ctga	cqq	ctac	acac	ta c	ttaa	gcaac	300
228	acct	ctga	ga c	gtcc	ctat	a cc	aggt	ccta	cta	aaaa	caa	aaca	acta	at a	cado	cggga	360
230	ccac	acac	ta t	gtat	adda	a aa	tgag	acaa	ata	gaga	aca	3500	acta	ta a	aage	gcacg	420
232	geet	ccag	ca c	tgac	ataa	2 22	taat	3003	ata	gaga	gea	accc.	7000	ta c	cayy	geacg	
234	atcc	taca	-5 C	ataa	2523	a +~	-99-	2525	at a	9499	-ac	cagi	30001	uu Ci	acca.	attac	480
234	atas	at ~~	oy L	9-90	arge	u ug	accc above	uleg	gcg.	atCt'	Lüg	agac	agge	at ga	aact	gctgg	540
220	guda	t accord	uu g	gggc	agcc	u ca	ytga	ygaa	gac	ctcc.	tgc	ccga	accg	cg ga	atcc	tgcag	600
230	aaac	rcgci	ig t	gccc	atca	t cg	acaca	accc	aag	tgca	acc	tgct	ctaca	ag ca	aaag	acacc	660
240	gagt	ttgg	ct a	ccaa	ccca	a aa	ccat	caag	aat	gaca	tgc	tgtg	gcc	gg ct	ttcg	aggag	720
242	ggca	agaa	gg a	tgcc	tgca	a gg	gcga	ctcg	ggc	ggcc	CCC	tggt	gtgc	et e	ataa	gtcag	780
244	tcgt	ggct	gc a	ggcg	ggggi	t gat	cago	ctgg	ggtgagggct			gtgcccgcca qaaccqccca				840	
246	ggtg	tctad	ca t	ccgt	gtca	c cg	ccca	ccac	aact	tggat	taa .	atcq	atca	at co	ccca	aactg	900
248	cagt	tccad	gc c	agcq	aggti	t gad	gcaad	caq	aaqt	tctac	gac a	atcad	cate	a co	catica	actag	960
	-			-				_		-	-						

PATENT APPLICATION: US/10/040,655A

DATE: 11/17/2004 TIME: 12:29:19

```
250 cggccgcttc cctttagtga gggttaatgc ttcgagcaga catgataaga tacattgatg
                                                                         1020
252 agtttggaca aaccacaact agaatgcagt gaaaaaaatg ctttatttgt gaaatttgtg
                                                                         1080
254 atgctattgc tttatttgta accattataa gctgcaataa acaagttgac
                                                                         1130
257 <210> SEQ ID NO: 9
258 <211> LENGTH: 315
259 <212> TYPE: PRT
260 <213> ORGANISM: Artificial
262 <220> FEATURE:
263 <223> OTHER INFORMATION: PFEK-PROTT-HIS fusion protein amino acid sequence
265 <400> SEQUENCE: 9
267 Met Asp Ser Lys Gly Ser Ser Gln Lys Ser Arg Leu Leu Leu Leu
268 1
271 Val Val Ser Asn Leu Leu Cys Gln Gly Val Val Ser Asp Tyr Lys
272
                20
                                    25
275 Asp Asp Asp Val Asp Ala Ala Leu Ala Ala Pro Phe Asp Asp
                                40
279 Asp Asp Lys Ile Val Gly Gly Tyr Ala Leu Glu Glu Gly Glu Trp Pro
283 Trp Gln Val Ser Ile Gln Arg Asn Gly Ser His Phe Cys Gly Gly Ser
                        70
287 Leu Ile Ala Glu Gln Trp Val Leu Thr Ala Ala His Cys Phe Arg Asn
                    85
291 Thr Ser Glu Thr Ser Leu Tyr Gln Val Leu Leu Gly Ala Arg Gln Leu
                                    105
295 Val Gln Pro Gly Pro His Ala Met Tyr Ala Arg Val Arg Gln Val Glu
296
                                120
                                                    125
299 Ser Asn Pro Leu Tyr Gln Gly Thr Ala Ser Ser Ala Asp Val Ala Leu
        130
                            135
303 Val Glu Leu Glu Ala Pro Val Pro Phe Thr Asn Tyr Ile Leu Pro Val
304 145
                        150
                                            155
307 Cys Leu Pro Asp Pro Ser Val Ile Phe Glu Thr Gly Met Asn Cys Trp
                    165
                                        170
311 Val Thr Gly Trp Gly Ser Pro Ser Glu Glu Asp Leu Leu Pro Glu Pro
312
                180
                                    185
315 Arg Ile Leu Gln Lys Leu Ala Val Pro Ile Ile Asp Thr Pro Lys Cys
           195
                                200
319 Asn Leu Leu Tyr Ser Lys Asp Thr Glu Phe Gly Tyr Gln Pro Lys Thr
                            215
323 Ile Lys Asn Asp Met Leu Cys Ala Gly Phe Glu Glu Gly Lys Lys Asp
                                            235
327 Ala Cys Lys Gly Asp Ser Gly Gly Pro Leu Val Cys Leu Val Gly Gln
                    245
                                        250
331 Ser Trp Leu Gln Ala Gly Val Ile Ser Trp Gly Glu Gly Cys Ala Arg
                                    265
335 Gln Asn Arg Pro Gly Val Tyr Ile Arg Val Thr Ala His His Asn Trp
           275
                               280
339 Ile His Arg Ile Ile Pro Lys Leu Gln Phe Gln Pro Ala Arg Leu Gly
                           295
343 Gly Gln Lys Ser Arg His His His His His
```

PATENT APPLICATION: US/10/040,655A

DATE: 11/17/2004 TIME: 12:29:19

```
344 305
                              310
                                                   315
     347 <210> SEQ ID NO: 10
     348 <211> LENGTH: 4
     349 <212> TYPE: PRT
     350 <213> ORGANISM: Artificial
     352 <220> FEATURE:
     353 <223> OTHER INFORMATION: Chromogenic substrate
     356 <220> FEATURE:
     357 <221> NAME/KEY: MISC_FEATURE
     358 <222> LOCATION: (1)..(1)
     359 <223> OTHER INFORMATION: N-Succinyl-alanine
     361 <220> FEATURE:
     362 <221> NAME/KEY: MISC_FEATURE
     363 <222> LOCATION: (4)..(4)
     364 <223> OTHER INFORMATION: Phe-p-nitroanilide
     366 <400> SEQUENCE: 10
W--> 368 Xaa Ala Pro Xaa
     369 1
     372 <210> SEQ ID NO: 11
     373 <211> LENGTH: 4
     374 <212> TYPE: PRT
     375 <213> ORGANISM: Artificial
     377 <220> FEATURE:
     378 <223> OTHER INFORMATION: Chromogenic substrate 6
     381 <220> FEATURE:
     382 <221> NAME/KEY: MISC FEATURE
     383 <222> LOCATION: (1)..(1)
     384 <223> OTHER INFORMATION: N-(Methoxysuccinyl)-Ala
     386 <220> FEATURE:
     387 <221> NAME/KEY: MISC FEATURE
     388 <222> LOCATION: (4)..(4)
     389 <223> OTHER INFORMATION: Phe-p-nitroanilide
     391 <400> SEQUENCE: 11
W--> 393 Xaa Ala Ala Xaa
     394 1
```

RAW SEQUENCE LISTING ERROR SUMMARY PATENT APPLICATION: US/10/040,655A

DATE: 11/17/2004 TIME: 12:29:21

Input Set : A:\ORT1566NP.Subst.Seq.List.txt
Output Set: N:\CRF4\11172004\J040655A.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:10; Xaa Pos. 1,4 (Seq#:11; Xaa Pos. 1,4

Invalid <213> Response:

Use of "Artificial" only as "<213> Organism" response is incomplete, per 1.823(b) of New Sequence Rules. Valid response is Artificial Sequence.

Seq#:2,3,4,5,6,8,9,10,11

VERIFICATION SUMMARY

PATENT APPLICATION: US/10/040,655A

DATE: 11/17/2004 TIME: 12:29:21

Input Set : A:\ORT1566NP.Subst.Seq.List.txt
Output Set: N:\CRF4\11172004\J040655A.raw

L:368 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10 after pos.:0 L:393 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11 after pos.:0